

1 atgggtcacg gtgtcagctg cgcccgacacc ggcgacgagc acgacttctt ccggggcg
61 cagctggcg acctcgacgc cctggccgccc ctccctcgccg ccgaccccttc cctcgctcg
121 cgcgcaccc tctacgacccg cctctccgtc ctccacatcg ccgcccggaa tggccgcata
181 gaggtgtctt ccatgttctt ggatcgccgg ggcgcggccgg acgcggtaa tcggcacaag
241 cagacgcgc tgatgctcgc ggcacatgcac ggcaagatcg actgcgtgct caagctctc
301 caggccgacg caaatatctt gatgttcgac tcgggtcacg cgaggacctg cttccaccac
361 gcggcgtact acggccacgt cgactgcctg caggccatcc tcggccggc gcagaccacg
421 ccggggccg actcatgggg ttgcggccgg ttgcgtcaacg tcagggacga ccacggcgcc
481 actccgcgtc atctcgccgc caggcagggg cggccgggggt gcgtgcagggt ttgtggag
541 aacggcgcca ttgtgtcgcc ttgacagga tcatatggct tccctggaa cacgtcgctt
601 catggctg ctcgttagcgga gaactggat tgcatcgga agctgtgc ctggggagct
661 gatcggtcc aaagggttc ggctggaga attccctatt ctgtgtcgct gaaacggAAC
721 catggagcat gtgcagctt gctgaacctt acatcagcag agccccatggt gtggccatcc
781 ccactaagt tcatcagtga gcttgaacca gaagctaagg ctctcctgga agcagcttg
841 atggaaagcca acaggggagag ggagaagaaaa atcctgaatg gcacaaagta ctccctgcca
901 tcccttcgc cgggtgatga cagtggcat gacgatgcat gctcagaggt gagcgcacacg
961 gagcttgct gcatctgctt cgaccaggct tgccattt aggtgcaga ctgtggacat
1021 caaatgttg caccgtgcac gctggactg tgctgtcaca acaaacccaa tccgacgacc
1081 ctgacaccgc cctcaccggc ctggccatcc tgccggggca gcatctcagc gctgggttg
1141 gccaaaacaa ggctgtctt tgatcctgac aagccgtcat ccctgcagct cacccggaaag
1201 cggtcgcgtc gatctcacaa cctcagttag ggcagcagca gcttcaaagg gtcacccctcg
1261 gccatgggct ccttctcaaa gcttggccgt ggctcgagcc gcatggcgga cagtgcacagc
1321 agcaaccttgg acaaggctga gcacgatcta tga

FIG. 1A

I	MGHGVSCART	10
II	GDEHD FFRAAHLD ALAALLAADPSLARRATLY DRLSV LHIAANGRIEVLSMFL DRGAPPDAVNR HKQTPL MLAAMHGKIDCVLKLLQADANILMFDSV HARTC LHAAYYGHVDCLQAILAAAQTTPVADSWG FARFVNVRDD HGATPL LHAAARQGRPGCVQVILLENGAIVSALTGSYGF PGSTS LHAAARSGNLD CIRKLLAWGADRLQRDSAGRIPYSVA LKRNHGAC AALLNPTSAEPMVWPSPLKFISELE PEAK ALLEALMEANREREKKILNGTKYSLPSPSPG	45 78 112 157 194 236 269 305
III	DDSADDDACSEVS	318
IV	DTELCCICFDQACTIEVQDCG RQMCA PCTLALCCHNKPNPTTLTP PSPACPF ORGSI SRLVVAQTRS	363 385
V	ACDPDKPSSLQLTRKRSRRSHNLSEGSSFKGLPSAMGSFSKLGR GSSRMADSDSSNLDKPEHDL	430 450

FIG. 1B

XB3
c-Cbl
IAP

CCICFDQACTIEVQDCGHQM-CAPCTLALCCHNKPNTTLPPSPACPFCR
CKICAENDKDVKIEPCGHLM-CTSCLTSWQESEGQG-----CPFCR
CKICYVEECIVCFVPCGHVVACAKCALSV-----DKCPMCR

FIG. 1C

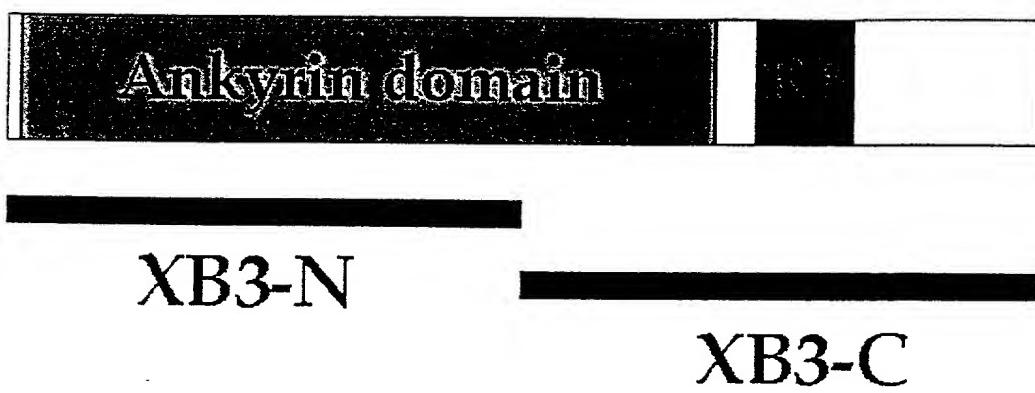


FIG. 2